

EXHIBIT B

A Comprehensive Guide for the Accurate Classification of Murine Hair Follicles in Distinct Hair Cycle Stages

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Numerous strains of mice with defined mutations display pronounced abnormalities of hair follicle cycling, even in the absence of overt alterations of the skin and hair phenotype; however, in order to recognize even subtle, hair cycle-related abnormalities, it is critically important to be able to determine accurately and classify the major stages of the normal murine hair cycle. In this comprehensive guide, we present pragmatic basic and auxiliary criteria for recognizing key stages of hair follicle growth (anagen), regression (catagen) and quiescence (telogen) in C57BL/6NCrIBR mice, which are largely based on previous work from other authors. For each stage, a schematic drawing and representative micrographs are provided in order to illustrate these criteria. The basic criteria can be employed for all mouse strains and require only routine histochemical techniques. The auxiliary criteria depend on the immunohistochemical analysis of three markers

(interleukin-1 receptor type I, transforming growth factor- β receptor type II, and neural cell-adhesion molecule), which allow a refined analysis of anatomical hair follicle compartments during all hair cycle stages. In contrast to prior staging systems, we suggest dividing anagen III into three distinct substages, based on morphologic differences, onset and progression of melanogenesis, and the position of the dermal papilla in the subcutis. The computer-generated schematic representations of each stage are presented with the aim of standardizing reports on follicular gene and protein expression patterns. This guide should become a useful tool when screening new mouse mutants or mice treated with pharmaceuticals for discrete morphologic abnormalities of hair follicle cycling in a highly reproducible, easily applicable, and quantifiable manner. **Key words:** alkaline phosphatase/anagen/catagen/dermal papilla/telogen. *J Invest Dermatol* 117:3–15, 2001

The hair follicle (HF) is a highly sensitive mini-organ whose cyclic transformations from phases of rapid growth (anagen), via apoptosis-driven regression (catagen) to relative quiescence (telogen) (Dry, 1926) are profoundly influenced by numerous growth factors, cytokines, hormones, neuropeptides, and pharmaceutical products (for review see Paus, 1996, 1998; Stenn *et al*, 1996, 1998; Paus and Cotsarelis, 1999). These manipulations of HF cycling are of substantial interest to an ever-growing community of life scientists and physicians. Yet, it is far from trivial to assess accurately and quantify alterations in HF cycling.

In addition to the human hair cycle (Kligman, 1959), the main three phases of the HF cycle have been described and investigated in most pigmented mouse strains, albino mice, and in some mouse mutants (Dry, 1926; Chase *et al*, 1951; Chase, 1954; Orwin *et al*,

1967; Sundberg, 1994; Panteleyev *et al*, 1998, 1999). Decades ago, landmark publications by Chase *et al* (1951), Chase (1954), and Straile *et al* (1961) defined key parameters for the recognition of distinct stages of the murine hair cycle. For nearly 50 y, these publications have been used as key references for HF classification, although none of them offers a comprehensive, unified classification scheme for the complete hair cycle; however, describing even subtle hair cycle changes has become of paramount importance to the analysis of mouse mutants, transgenic and knockout mice, and to the analysis of experimental drug effects in mice. The present guide aims to offer scientists with an interest in hair research an updated and pragmatic approach to rapid, instructive comparative analyses of murine hair growth patterns. This guide complements our earlier guide on the classification of murine HF development (Paus *et al*, 1999), which combined provide a standardized method for the analysis of murine HF growth.

Based on fundamental histologic and ultrastructural studies on murine cycling HF (Dry, 1926; Butcher, 1951; Chase *et al*, 1951; Wolbach, 1951; Straile *et al*, 1961; Parakkal, 1969a, b, c, 1970; De Weert *et al*, 1982), we have summarized pragmatic criteria for the recognition of distinct stages of the hair cycle (Figs 1 and 2).

This guide suggests basic as well as more advanced auxiliary criteria to define anagen, catagen, and telogen stages of the hair cycle (Figs 1 and 2), which are widely applicable to different

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Abbreviations: a.d., after depilation; AP, alkaline phosphatase; APM, arrector pili muscle; CTS, connective tissue sheath; DP, dermal papilla; ES, epithelial strand; HF, hair follicle(s); HS, hair shaft; NCAM, neural cell-adhesion molecule; SG, sebaceous gland.

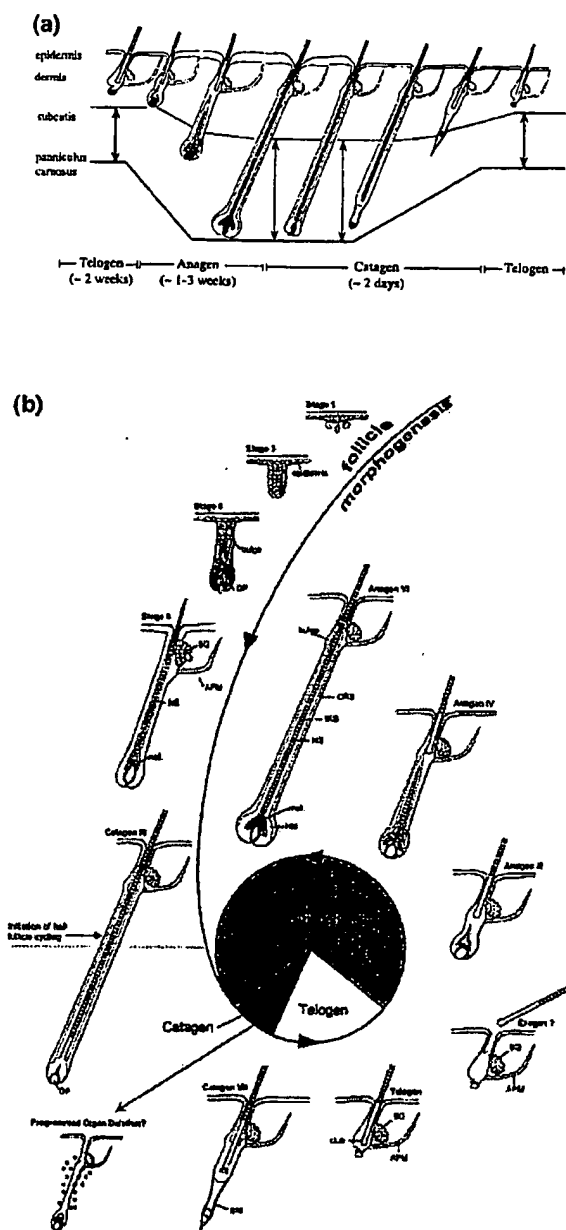


Figure 1. (a) Schematic representation of the increasing and decreasing length of the HF and the localization of the most proximal part of the HF in correlation with the panniculus carnosus and the border between the dermis and subcutis. Arrows between panniculus carnosus and the border dermis/subcutis indicate the hair cycle-associated changes in the thickness of the subcutis. The approximate duration of each phase is indicated in brackets. Note changes of the DP shape and size throughout the cycle as well as the increasing size of the SG during anagen IV–VI. (b) Schematic representation of key stages of HF development and cycling. Stages 1–8 represent stages of HF development. Anagen: growth phase, catagen: regression phase, telogen: “resting” phase, exogen: (speculative) active shedding phase. For detailed description of programmed organ deletion see Eichmüller *et al.*, 1998, for exogen see Stenn and Paus, 2001; black dots refer to immune cells. APM, arrector pili muscle; BM, basement membrane; DP, dermal papilla; HS, hair shaft; IRS, inner root sheath; mel, melanin; ORS, outer root sheath; SG, sebaceous gland.

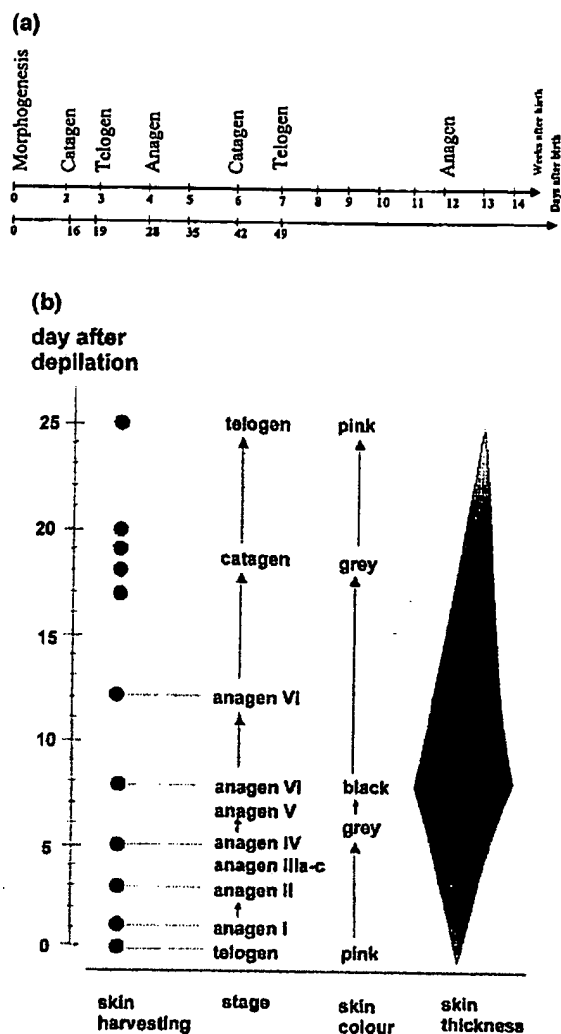


Figure 2. (a) Time-scale for the hair cycle in female C57BL/6 mice during the first 14 wk after birth. (b) Time-scale for the hair cycle in female C57BL/6 mice and changes of skin pigmentation and skin thickness after depilation.

mouse strains and mutants. In essence, this classification guide can also be utilized for staging the HF of other hairy animals, even though species-specific anatomic differences must be taken into account. The staging guide for mice is also highly relevant for human hair biology, as the same basic follicle transformations occur in mice and men (Kligman, 1959). Until a rigorous comparison of the more subtle morphologic characteristics of murine and human HF cycling has been published, however, it is wise not to claim that the current guide can be used uncritically for human HF staging as well.

HF morphogenesis as well as the first postnatal catagen, telogen, and anagen development follow a rather precise time-scale (Paus *et al.*, 1999) (Fig 2a). Nevertheless, these processes are dependent on the genetic background (mouse strain), the sex (e.g., female mice show a prolonged telogen) as well as environmental factors such as time of the year (temperature, light periods) and nutritional factors. To avoid associated fluctuations, this current guide is based

on the highly standardized C57BL/6NCrIBR (C57BL/6) model of depilation-induced HF cycling (Fig 2b) (Chase, 1954; Paus *et al*, 1990, 1994a, b). Briefly, a wax/rosin mixture is applied on the dorsal skin of 7 wk old mice with all dorsal skin HF in telogen, as evidenced by the homogeneous pink skin color. Removing the wax/rosin mixture removes all hair shafts and immediately induces homogeneous anagen development over the entire depilated back of the mouse. After full anagen development, the consecutive stages (catagen and telogen) are then entered spontaneously in a fairly homogeneous manner. Compared with spontaneous anagen development, there are two major differences: (i) depilation-induced anagen is fully synchronized over the entire area of depilation, whereas spontaneous anagen develops in a wave-like pattern, and (ii) a slight inflammatory effect of plucking ("wounding response") has been demonstrated (Argyris, 1968), which can also be appreciated by upregulation of epidermal intercellular adhesion molecule-1 immunoreactivity 1 d after depilation (Müller-Röver *et al*, 2000a). Similar to spontaneous anagen, even anagen induced by cyclosporine A, a potent immunosuppressant drug, does not show significant morphological differences compared with depilation-induced hair cycling (Paus *et al*, 1998). No significant differences in the expression patterns of interleukin (IL)-1 α , transforming growth factor (TGF)- β RII or neural cell-adhesion molecule (NCAM) have been found in these three different models (Paus and Müller-Röver, unpublished observation).

Owing to the strict coupling of follicular melanogenesis and HF cycling, anagen development is associated with characteristic changes in skin pigmentation (Fig 2b) (Slominski *et al*, 1991, 1994; Slominski and Paus, 1993). In addition, synchronized HF cycling in mice induces profound alterations in the architecture and thickness of almost all skin compartments, which are most evident in substantial, hair cycle-associated fluctuations in skin thickness (Fig 2b) (Chase, 1954; Paus *et al*, 1990, 1991). Nine days after depilation, the induced anagen HF reach their maximal length and are morphologically indistinguishable from spontaneously developing anagen follicles, even though the plucking trauma induces a short wound healing response immediately after the depilation (Chase *et al*, 1951; Silver *et al*, 1969; Slominski *et al*, 1991; Paus *et al*, 1998). In contrast to the normal caudal-nuchal development of a spontaneously developing anagen wave, spontaneous catagen development starts in the neck region and proceeds in a nuchal-

caudal direction. In C57BL/6NCrIBR mice, catagen-associated changes in HF morphology are first seen on day 17 after depilation in the neck region, and are macroscopically recognizable by a switch in skin color from black to gray-pink. The catagen wave reaches the tail region about 2 d later on day 19/20 (Chase *et al*, 1951; Chase, 1954; Straile *et al*, 1961) (Fig 2).

HOW TO USE THIS GUIDE

Although spontaneous HF cycling is envisioned to begin with catagen (Paus and Cotsarelis, 1999; Stenn, 1999; Stenn and Paus, 2001) (Fig 2a), hair cycle studies customarily begin with the telogen-anagen transformation. This is reflected in the current guide (Fig 3). In order to avoid terminologic confusion, which is often caused by differences in how selected terms are used in hair research papers, we have summarized definitions of key terms employed in the context of this guide (Table I). Please note that the term "proximal" here refers to those parts of the HF that are located close to the panniculus carnosus, whereas "distal" refers to those parts located close to the epidermis. In order to illustrate key parameters for the recognition of distinct HF stages throughout the hair cycle, Fig 3 is structured as follows: the left-hand column shows computer-generated schematic drawings of six distinct main stages and three substages of HF growth (anagen), eight stages of HF regression (catagen) and the quiescence stage (telogen) of the hair cycle, modified from previous works (Chase *et al*, 1951; Straile *et al*, 1961; Paus *et al*, 1997) (for explanation and definition of the corresponding terminology, see Table I). The criteria presented here are highly reproducible and reliable in all C57/BL6 mice over a very wide age range, which have been investigated in our group during the last 10 y. These criteria have been described in most pigmented mouse strains, albino mice, and in some mouse mutants (Dry, 1926; Chase *et al*, 1951; Chase, 1954; Orwin *et al*, 1967; Sundberg, 1994; Panteleyev *et al*, 1998, 1999) and are valid also in later cycles of 1 y old mice (Paus and Müller-Röver, unpublished observation).

The central column in Fig 3 provides a list of basic and auxiliary classification criteria that are separated from each other by a dotted line (top: basic, bottom: auxiliary criteria). The former are recognizable by routine light microscopy, whereas the latter require additional staining methods. The basic criteria are applicable to all mouse strains and are not dependent on the presence of

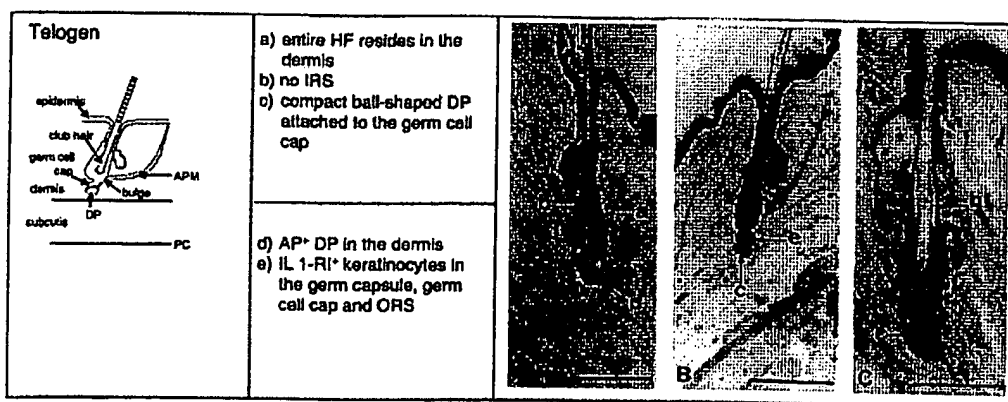


Figure 3. A comprehensive guide for the recognition and classification of distinct stages of hair cycling. The left-hand column shows a computer-generated schematic drawing of the so-called quiescence phase (telogen) of the hair cycle. The second column summarizes essential basic criteria for recognition of the single substages (above dotted line) and auxiliary criteria for more precise staging (below dotted line). The right-hand column shows representative micrographs of each hair cycle stage (lower case letters correspond to lower case letters used in the central column). The following staining techniques were employed: (A, C) Giemsa (Romeis, 1991); (B) IL-1 RI immunoreactivity. (Eichmüller *et al*, 1998). Scale bars: 50 μ m. Please note: upper case letters in the left-hand corner label the image whereas lower case letters identify tissues corresponding to the lower case letters of the criteria listed in the central column. Please also note: the left-hand column lists comprehensively all helpful markers, although not all markers are shown in the micrographs.

Table I. Glossary of anatomical terms frequently used in hair research^a

Term	Definition
Bulb	Prominent, onion-shaped thickening on the proximal end of the HF; consists of relatively undifferentiated matrix cells, HF melanocytes and of cells from the proximal ORS
Bulge	Convex extension of the distal part of the ORS, near the epidermis, location of epithelial follicle stem cells and point of insertion of the <i>m. arrector pili</i> .
Club hair	Residing hair shaft with a hollow brush of keratinized keratinocytes on the proximal end, tightly attached to the cortical cells of the hair cortex
Connective tissue sheath (CTS)	Part of the dermal connective tissue, tightly attached to the outer side of HF, composed of fibroblasts (and macrophages) and connective tissue
Dermal papilla (DP) syn: follicular papilla (FP)	Mesodermal part of the HF, which consists of closely packed mesenchymal cells; framed by the bulb matrix during anagen
Epithelial strand (ES)	Column of epithelial cells between the germ capsule and the compact DP; laterally demarcated by the thickened glassy membrane
Secondary germ capsule syn: secondary hair germ	Bag-like structure of glycogen-free cells (germ cells) of distal ORS, surrounding the club hair
Hair shaft	The hair <i>per se</i> , composed of trichocytes (= terminally differentiated HF keratinocytes), divided into hair cuticle, cortex and medulla
Hair shaft medulla	Central part of the hair shaft, composed of large, loosely connected keratinized cells with large intercellular air spaces
Hair shaft cortex	The mass of the hair shaft, composed of keratinized cells, longitudinally packed with keratin filaments (and melanin granules in pigmented hair shafts)
Hair canal	Passage way between epidermal surface and the most distal part of the IRS, demarcated by surrounding ORS.
Hyaline membrane syn: vitreous membrane, glassy membrane	Outermost noncellular part of the HF; composed of basal lamina and two layers of orthogonally arranged collagen fibers; separates ORS from CTS
Infundibulum	Most distal part of the HF in the dermis extending from sebaceous duct to the epidermal surface (including hair canal and distal ORS)
Isthmus	Middle portion of the HF extending from the sebaceous duct to the insertion of the <i>m. arrector pili</i> (bulge region)
Inner root sheath	Multilayered structure composed of terminally differentiated HF keratinocytes surrounded by the ORS; consists of Henle's layer, Huxley's layer and cuticle; surrounds the hair shaft up to the hair canal(IRS)
Outer root sheath (ORS)	Outermost sheath of HF keratinocytes, which merges distally into the basal layer of the epidermis and proximally into the hair bulb
Papillary stock syn: papillary stalk	Fibroblasts that link the proximal pole of the DP and the perifollicular CTS
Sebaceous gland	Glandular structure close to the insertion of the <i>m. arrector pili</i> with holocrine function; composed of lipid-filled sebocytes with a foamy appearance

^aFor introductory references, see Chase *et al* (1951); Wolbach (1951); Montagna and Ellis (1958); Montagna and van Scott (1958); Straile *et al* (1961); Roth (1965); Parakkal (1969a); Parakkal (1969c, b, 1970); De Weert *et al* (1982); Cotzarelis *et al* (1990); Abell (1994); Paus *et al*, 1994b; Paus *et al* (1998); Paus and Cotzarelis (1999); Stenn and Paus, 2001.

Table II. Synopsis of auxiliary methods used in this guide^a

Auxiliary methods/markers	Compartment	Stage	Reference
Pigmented mouse strain Melanin granules	Keratogenic region	Anagen IIIa-catagen III	Slominski <i>et al</i> (1991, 1994); Slominski and Paus (1993)
Histochemistry AP staining	DP	All stages	Handjiski <i>et al</i> (1994)
Oil-red-O staining	Sebocytes and hair canal	Anagen IIIc-V	Romeis (1991)
TUNEL staining	hair matrix, ORS, IRS	Catagen I-IV	Lindner <i>et al</i> (1997)
Immunohistochemistry IL-1 RI	ORS (in contrast to IRS)	All stages	Eichmüller <i>et al</i> (1998)
TGF- β RII	ORS (in contrast to IRS)	All stages	Paus <i>et al</i> (1997)
NCAM	DP, perifollicular CTS and CTS tail	Catagen IV-VIII	Müller-Röver <i>et al</i> (1998)
P-cadherin	Papillary cap, inner hair matrix	Telogen-anagen II	Müller-Röver <i>et al</i> (1999)

^aThese methods provide additional help to determine the stage-specific morphology of distinct HF compartments such as the DP or the papillary cap. Additionally, key stages are listed where these methods are of particular value in difficult cases.

pigment granules (melanin) in the HF. The auxiliary criteria that require a pigmented mouse strain such as C57BL/6J, and/or enzyme histochemical techniques, i.e., alkaline phosphatase (AP) and TUNEL staining; or immunohistochemistry for cytokine receptors, such as IL-1 receptor type I (IL-1 RI) and TGF- β receptor type II (TGF- β RII) and adhesion receptors, e.g. NCAM (Table II). The right-hand column in Fig 3 illustrates the criteria

listed in the central column with three representative micrographs. The lower case letters used for indicating key parameters correspond to the lower case letters in the central column list. Note that not all parameters are necessarily shown in the right-hand columns.

It is of important to note that the markers used in this guide are by no means the only useful markers for HF staging. For example, staining for soft and hard keratins likely is of great help for the

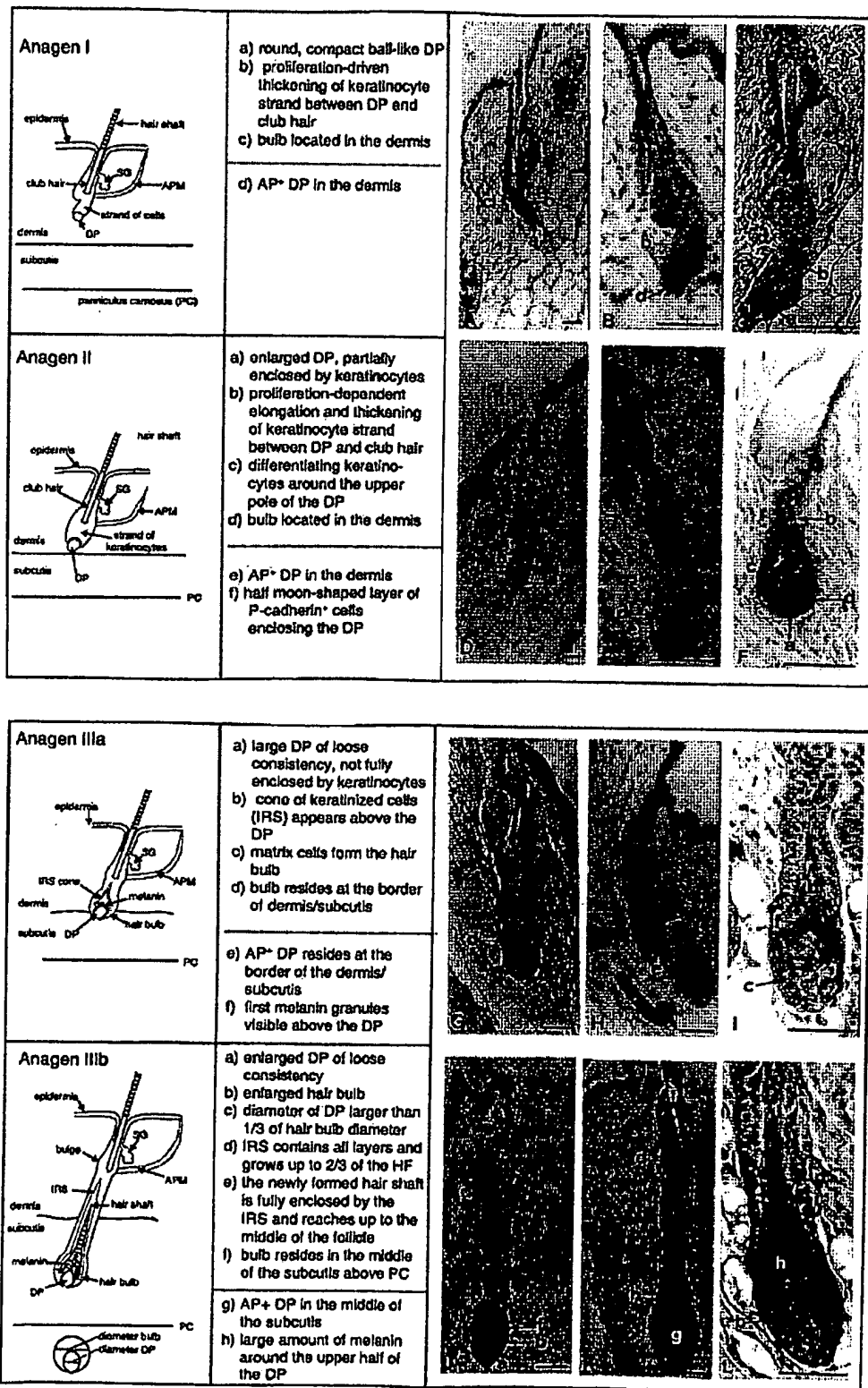


Figure 4. Legend on page 9.

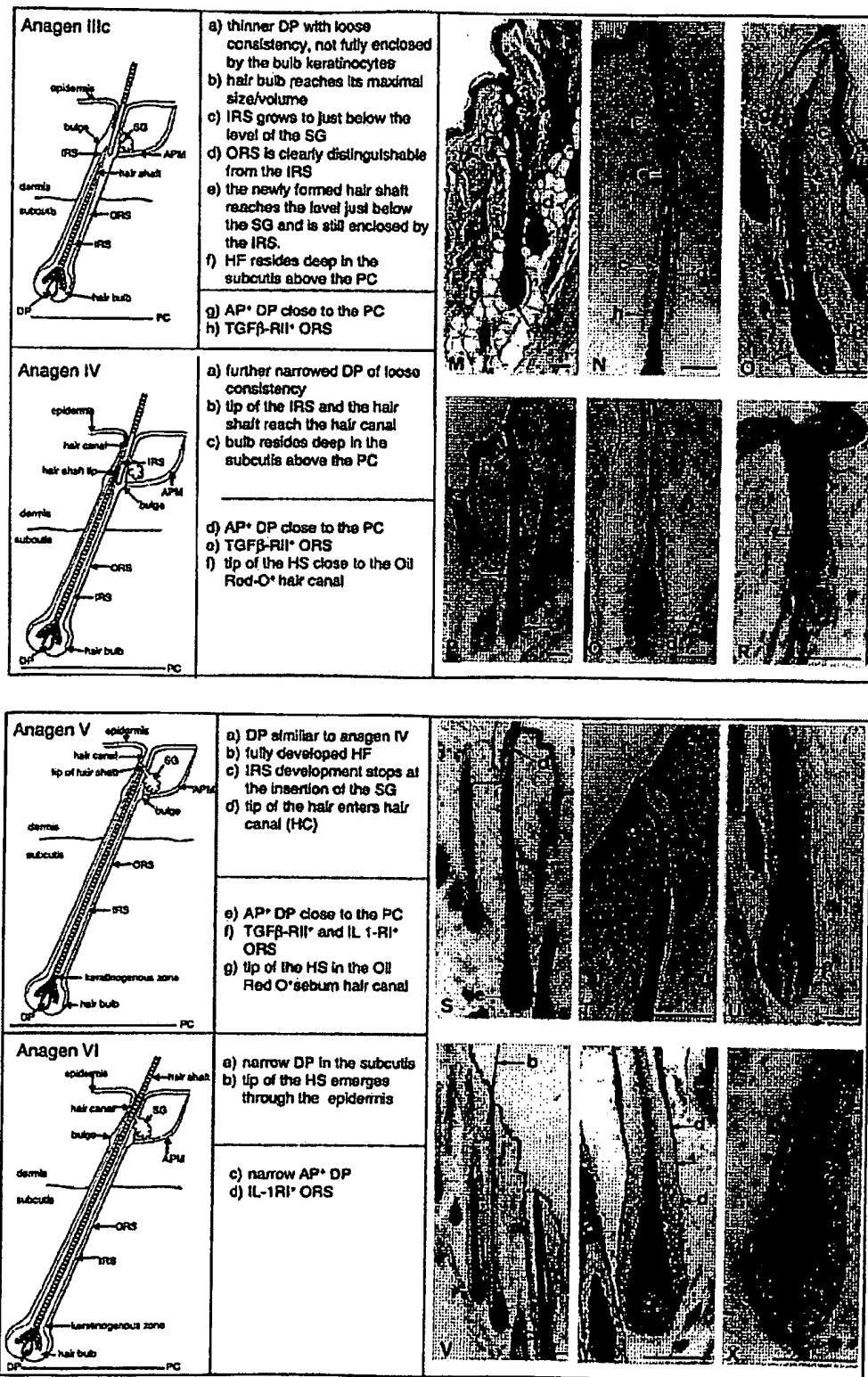


Figure 4. Continued.

recognition and accurate classification of murine HF in different hair cycle stages. Unfortunately, most anti-mouse keratin antibodies we have tested so far show massive background staining, and considerable variations of staining patterns are seen in specific anatomical compartments of the HF at distinct hair cycle stages (Müller-Röver and Paus, unpublished observation). Therefore, we have decided to use antibodies against IL-1R1, TGF- β RII, and NCAM for staining purposes, as these markers show very reliable, highly reproducible follicular staining patterns in specific anatomical compartments throughout hair cycling and experimental hair cycle modulations. Nevertheless, one always needs to consider whether pharmacologic or genetic manipulations in the animals that are to be analyzed might have changed the expression patterns of these markers independently of hair growth and regression (for example: NCAM upregulation in *Msx-2* overexpressing mice, Jiang *et al.*, 1999).

CRITERIA FOR THE RECOGNITION OF DISTINCT HAIR CYCLE STAGES

Stage-dependent HF length and skin thickness One of the first and easiest parameters used to classify the stage of a HF is its length (Fig 1b), which is measured from the dermal papilla (DP) to the epidermis. Anagen I–VI development (Fig 4) is characterized by increasing length of the HF and catagen I–VII by decreasing length (Fig 5). The HF reaches its maximal length during anagen VI (Fig 4V–X) and keeps its length during catagen I and II (Fig 5A–F). Thus, during anagen VI to catagen II, the DP is located close to the panniculus carnosus. During telogen, the HF reaches its minimal length (Fig 3). Because dermal fibroblasts surround the entire HF during anagens I and II (Fig 4A–F) it is strongly advised to use the AP staining method to visualize the DP. Easy to perform, the AP staining method takes only a few minutes, and prevents very common errors during HF staging such as mistaking the epithelial cells of the secondary hair germ of telogen or late catagen HF for fibroblasts of the DP.

As summarized in Figs 1(b) and 2(b), synchronized HF cycling in mice is also associated with stage-dependent changes of skin thickness and pigmentation. These independent parameters can be utilized to double-check the histomorphometric classification illustrated in Figs 3–5.

Basic rules for the recognition of telogen and anagen I and II HF A HF is in telogen, anagen I or anagen II as long as it is entirely surrounded by dermal fibroblasts and has not yet reached the subcutis. In addition, in pigmented mice, no melanin is visible in the hair matrix above the DP, although occasionally some

melanin granules from the preceding anagen phase can be observed in the DP, where they sometimes accumulate (rather than being extruded together with the hair shaft).

Stage-specific characteristics of telogen and anagen I and II HF Telogen HF (Fig 3 part 1A–C) are very easy to recognize as they are fully surrounded by interfollicular dermal fibroblasts (Fig 3B). They do not display any part of the inner root sheath (IRS) and the compact ball-shaped DP is closely attached to a small cap of secondary hair germ keratinocytes (Fig 3A, B). To distinguish DP and secondary hair germ, either the DP should be visualized by the AP staining method (Fig 3A) or the secondary hair germ at the upper pole of the DP should be marked by IL-1 R1 immunoreactivity (Fig 3B) or P-cadherin immunoreactivity (Müller-Röver *et al.*, 1999). In case the morphologic criteria described above are insufficient to distinguish telogen HF and late catagen HF, NCAM immunoreactivity can be employed as a key marker for making this distinction as NCAM immunoreactivity does not reveal more than one or two single NCAM⁺ connective tissue sheath (CTS) cells in close vicinity to the DP of telogen HF (not shown) whereas late catagen HF display a long tail of trailing NCAM⁺ CTS fibroblasts (Müller-Röver *et al.*, 1998). The number of club hairs in the hair canal is not a reliable parameter for HF staging. In late anagen at least two hair shafts fill the hair canal, the resting shaft from the preceding cycle and the new anagen VI hair shaft. At the end of telogen, some hair shafts are actively shed in a process termed exogen (Stenn *et al.*, 1998), whereas others may remain in the hair canal (Fig 4B). So far, no telogen substages have been proposed in the literature, but it is still controversially discussed whether the telogen phase is one homogeneous phase of resting (see Stenn *et al.*, 1998; Stenn and Paus, 2001).

Similar to telogen HF, anagen I HF (Fig 4A–C) are fully enclosed by the dermis. The only obvious difference to telogen HF (see below) is a thickening and prolongation of the strand of keratinocytes between the DP and the club hair. In contrast to anagen II HF, this keratinocyte strand has a smaller diameter than the secondary hair germ of the club hair. In difficult cases, P-cadherin staining is a helpful auxiliary method to visualize specifically the elongating strand of keratinocytes that cap the upper third of the round, compact ball-like DP (Müller-Röver *et al.*, 1999).

In contrast, anagen II HF (Fig 4D–F) are characterized by an enlarged DP compared with telogen or anagen I HF, which is more than half enclosed by proliferating keratinocytes of the developing hair matrix. The strand of keratinocytes between the DP and the old club hair, i.e., the developing hair bulb, now has a larger

Figure 4. A comprehensive guide for the recognition and classification of distinct stages of hair cycling. The left-hand column shows a computer-generated schematic drawing of the six distinct substages of hair growth (anagen) as suggested by Chase *et al.* (1951). The second column summarizes essential basic criteria for recognition of the single substages (above dotted line) and auxiliary criteria for more precise staging (below dotted line). The right-hand column shows representative micrographs of each hair cycle stage (lower case letters correspond to lower case letters used in the central column). The following staining techniques were employed: (A, C, D, F, G, J, M, O, T, U) Giemsa (Romeis, 1991); (W) IL-1 R1 immunoreactivity (Eichmüller *et al.*, 1998); (B, E, H, K, Q, S, V, X) AP staining (Handjiski *et al.*, 1994); (I, L, N) periodic acid-Schiff reaction (Romeis, 1991); (P) TGF- β RII immunoreactivity (Paus *et al.*, 1997); (R) Oil-Red-O labeling (Romeis, 1991). Scale bars: 50 μ m. Please note: upper case letters in the left-hand corner label the image whereas lower case letters identify tissues corresponding to the lower case letters of the criteria listed in the central column. Please also note: the left hand column lists comprehensively all helpful markers, although not all markers are shown in the micrographs.

Figure 5. A comprehensive guide for the recognition and classification of distinct stages of hair cycling. The left-hand column shows a computer-generated schematic drawing of the eight distinct stages of HF regression (catagen) as suggested by Straile *et al.* (1961). The second column summarizes essential basic criteria for recognition of the single substages (above dotted line) and auxiliary criteria for more precise staging (below dotted line). The right-hand column shows representative micrographs of each hair cycle stage (lower case letters correspond to lower case letters used in the central column). The following staining techniques were employed: (A, B, D, E, H, K, M, P, S, U, V, X) Giemsa (Romeis, 1991); (C, G, I, J, N, Q, T) AP staining (Handjiski *et al.*, 1994); (F, L) TUNEL (Lindner *et al.*, 1997); (O, R, W) NCAM immunoreactivity (Müller-Röver *et al.*, 1998). Scale bars: 50 μ m. Please note: upper case letters in the left-hand corner label the image whereas lower case letters identify tissues corresponding to the lower case letters of the criteria listed in the central column. Please also note: the left-hand column lists comprehensively all helpful markers, although not all markers are shown in the micrographs.

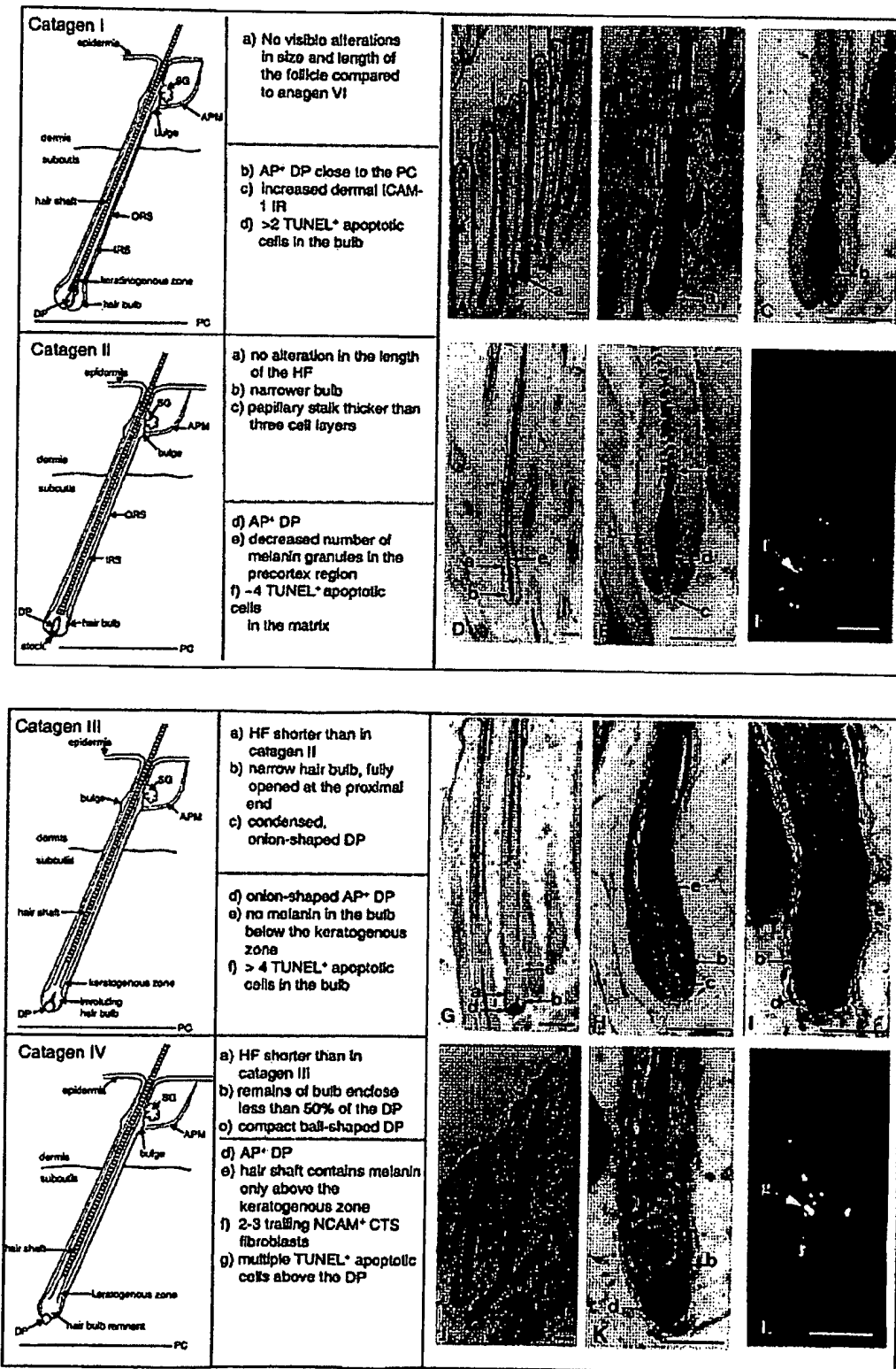


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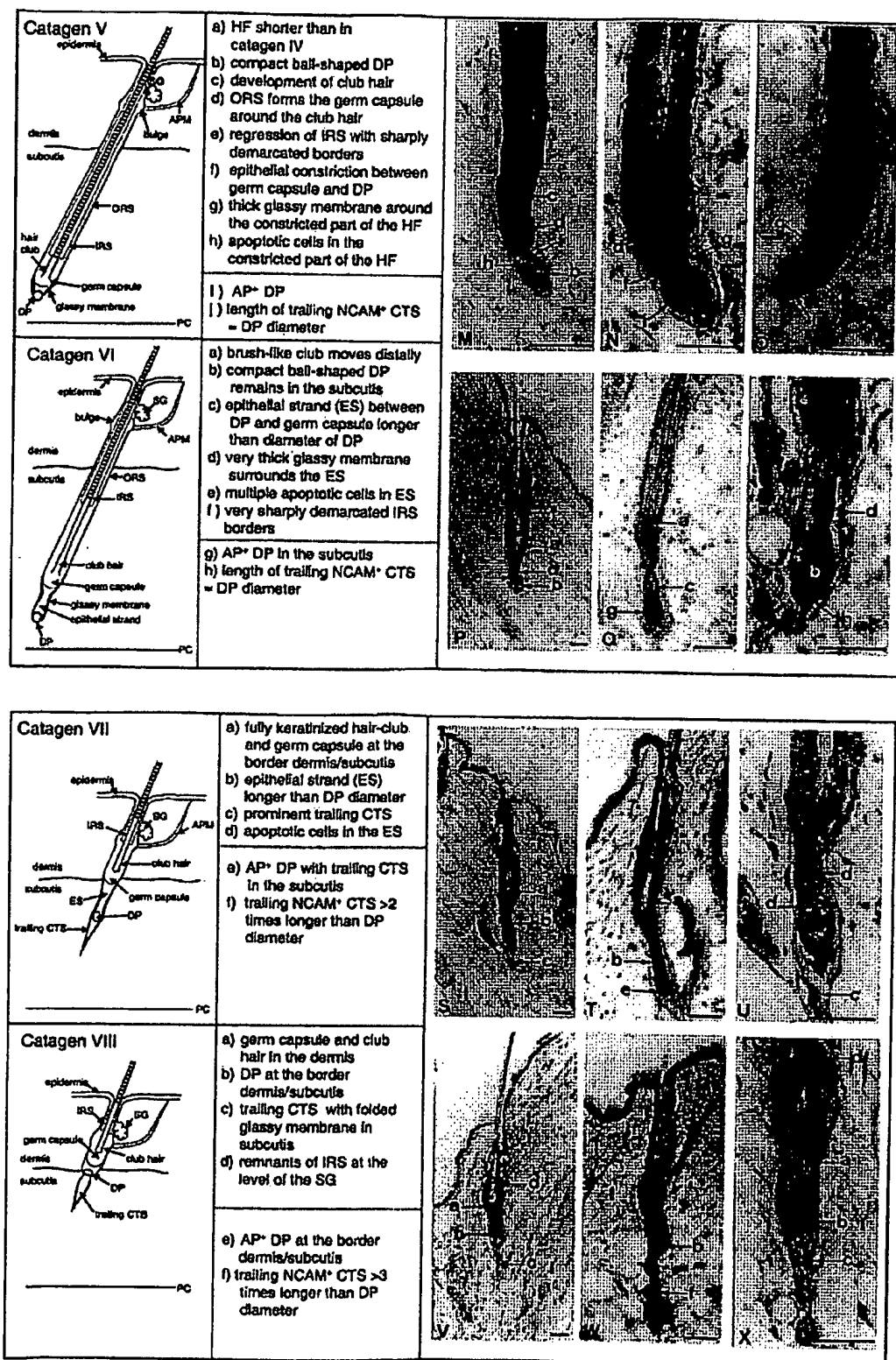


Figure 5. Continued.

diameter than the secondary hair germ of anagen I HF. The developing bulb and the DP are located at the border between the dermis and the subcutis. The upper pole of the DP of an anagen HF is surrounded by a half moon-shaped layer of keratinocytes that display a corona-like orientation, which one can easily visualize by P-cadherin immunoreactivity (Müller-Röver *et al*, 1999).

Anagen III: proposal to distinguish three distinct substages One aim of this guide is to standardize the description of hair cycle-dependent gene and protein expression patterns. In this context, it is a dilemma that the classical hair research literature (Chase *et al*, 1951; Chase, 1954; Straile *et al*, 1961) divides the hair cycle precisely in 15 different stages with a high descriptive value, whereas from a pragmatic point of view this may appear overdetailed and functionally insignificant. On the one hand, however, only this high degree of distinction allows one to detect even subtle abnormalities in HF cycling that had previously remained unnoticed (e.g., Maurer *et al*, 1997b; Botchkarev *et al*, 1998, 1999). On the other hand, it cannot be overemphasized that even "minor" alterations in follicle morphology from one of the well-defined substages of the hair cycle to the next reflect massive changes in gene expression, cell migration, proliferation, differentiation, apoptosis, and matrix remodeling. It is therefore conceivable that even more substages will have to be defined in the future, once the morphologic signs of cyclic HF transformation have been more rigorously and systematically lined up with the underlying changes in cell biology and gene expression.

The HF is in anagen III when the hair bulb surrounds the DP and resides at the border between dermis and subcutis or deeper, and when the tip of the newly formed IRS is below the hair canal. It is advisable to subdivide the historical stage "anagen III" into three substages for the following reasons. First, anagen III is the second longest phase of the murine hair cycle (approximately 3 d, with only anagen VI lasting longer; see Fig 2b) compared with the relatively short anagen I-II and IV-V phases and this encompasses massive morphologic changes in the follicle, which follow a easily distinguishable choreography of follicle transformation events. Secondly, anagen IIIa is characterized by the onset of melanogenesis (Chase *et al*, 1951; Slonimski *et al*, 1994) and the formation of the IRS cone, whereas anagen IIIb and IIIc are characterized by a fully developed IRS and a fully developed, pigmented hair shaft. Thirdly, the follicular DP is located at the border dermis/subcutis (in anagen IIIa), in the middle of the subcutis (in anagen IIIb) or close to the panniculus carnosus (in anagen IIIc). Finally, the bulb reaches its maximal size not before anagen IIIc (Chase *et al*, 1951; Chase, 1954). Each of these distinct processes is a primary target of pharmacologic or genetic manipulation and the effect of a given test agent may well differ between each of these three distinct substages of anagen III. Considering the magnitude of changes on the cell and extracellular matrix level (proliferation, migration, apoptosis, differentiation, melanogenesis, matrix synthesis) that corresponds to each of these anagen III substages, even this additional subdivision, in the future, may still be considered too crude. In the interest of practicability, however, it offers a reasonable compromise between user-friendliness and hair biologic necessities.

Characteristics of anagen IIIa, IIIb, and IIIc HF In anagen IIIa HF (Fig 4G-I) the bulb and DP reside at the border between the dermis and subcutis. During anagen IIIb (Fig 4J-L), the DP and bulb reside in the middle of the subcutis. Finally, during anagen IIIc (Fig 4M-O), the DP and bulb reside close to the panniculus carnosus.

Although in both cases the bulb is located close to the border between dermis and subcutis anagen II and IIIa HF are distinguished by the fact that anagen IIIa HF show the IRS cone above the DP and (in pigmented mice) melanin granules first appear within the developing IRS cone (Fig 4G-I). The developing IRS becomes detectable as a layer of differentiated keratinocytes arising from the most distal hair matrix and is visible as a cone-shaped structure that reaches up to the middle of the developing anagen

HF (Fig 4I). No hair shaft is visible at this time, as it forms only after the completion of anagen IIIa.

A developing hair shaft becomes clearly visible in anagen IIIb HF (Fig 4J-L). It reaches up to the middle of the HF, and is entirely surrounded by the IRS, which now contains all three of its layers (Henle's layer, Huxley's layer, cuticle). The distal tip of the IRS reaches up to two-thirds of the length of the HF, but still resides below the level of the old club hair, the insertion of the arrector pili muscle, and the sebaceous gland (SG) (Fig 4J-L). The bulb is now larger than in anagen IIIa, and fully encloses the enlarged DP. The DP now has a looser consistency and it makes up more than one-third of the bulb volume (see small insert diagram on p. 7 below the schematic representation of anagen IIIb). The bulb and DP of anagen IIIb HF reside in the mid-subcutis (Fig 4J-L).

During anagen IIIc (Fig 4M-O), the bulb and DP reach their deepest (i.e., most proximal) position and come to rest close to the panniculus carnosus in the subcutis (Fig 4M). The IRS and the hair shaft now reside just below the level of the SG, but do not reach the hair canal (Fig 4N, O). The outer root sheath (ORS) and IRS are clearly distinguishable (Fig 4N). In order to determine the precise location of the IRS tip, TGF- β RII immunoreactivity is a very useful auxiliary parameter because only the ORS is positive for this marker (Paus *et al*, 1997). Compared with anagen IIIb, the DP is now thinner and more elongated, but is still of loose consistency and not yet fully surrounded by hair bulb keratinocytes (Fig 4O).

Basic rule for the differentiation of anagen IIIc to anagen VI/catagen I HF The key technique for distinguishing anagen IIIc, IV, V and VI HF is to assess the exact position of the hair shaft and the IRS: during anagen IIIc, the hair shaft and IRS reach a level just below the insertion of the SG; during anagen IV, both hair shaft and IRS have reached the hair canal; during anagen V, the tip of the newly formed hair shaft enters into the hair canal; and during anagen VI, the tip of the hair shaft (early anagen VI) and beyond (late anagen VI/catagen I) emerge through the pilary canal to the skin surface.

Stage-specific characteristics of anagen IV to anagen VI/catagen I HF During the short anagen IV phase (Fig 4P-R), the bulb and DP still reside in the deep subcutis (Fig 4P, Q). Compared with anagen III, the bulb is enlarged and the DP is narrowed, thus, the diameter of the DP is smaller than one-third of the bulb diameter. That the distal end of the IRS and the tip of the hair shaft are closely attached to the hair canal (Fig 4R) serves as the basic parameter to distinguish anagen IV from the previous and consecutive stages. TGF- β RII immunoreactivity in the ORS gives a perfect outline of the TGF- β RII-negative IRS (Fig 4P). In order to determine precisely whether the hair shaft has already entered the hair canal, the Oil-Red-O technique (Romeis, 1991) is particularly helpful as it stains the sebum in the hair canal (Fig 4R).

Similar to the preceding stage, anagen V (Fig 4S-U) is a very short phase. Anagen V HF are easily recognized as at this time the tip of the hair shaft has entered the hair canal and is still fully enclosed by the ORS (Fig 4T). The DP has a similar shape, volume, and appearance as in anagen IV and VI (Fig 4S, U). IRS degradation begins at the level of the insertion of the SG (Fig 4T). In difficult cases, Oil Red-O staining helps to reveal the precise position of the tip of the hair shaft in the sebum-filled hair canal (cf. Fig 4R). When the tip of the hair shaft emerges through the epidermis, the HF enters anagen VI (Fig 4V). The narrow, elongated DP and the bulb reside close to the panniculus carnosus.

Anagen VI (Fig 4V, X) and catagen I HF (Fig 5A-C) cannot be distinguished by light microscopic morphologic criteria. For catagen, we use the TUNEL method to identify apoptotic keratinocytes. Any HF with more than two TUNEL⁺ cells in the bulb is defined as a catagen I HF, as the rate of TUNEL labeling is closely related to the morphologic changes during catagen development (Weedon and Strutton, 1984; Lindner *et al*, 1997).

Recognition of catagen II-IV HF The basic parameter characterizing catagen II-IV HF are: (i) the shape of the DP [as

it is best visualized by AP histochemistry (Handjiski *et al.*, 1994) or NCAM immunohistochemistry (Müller-Röver *et al.*, 1998)], and (ii) the shape of the bulb (catagen II HF displays a smaller epithelial bulb, which completely surrounds a long, oval DP; catagen III HF displays an onion-shaped DP that is predominantly, but not completely enclosed by the hair bulb; and catagen IV HF display a compact, ball-shaped DP that is surrounded by the bulb for less than half of its volume).

During catagen II (Fig 5D–F), the follicle retains its anagen VI/catagen I length (Fig 5D), but compared with anagen VI/catagen I HF has a smaller bulb (Fig 5D–F) and a narrower DP (Fig 5E, F). In addition, the papillary stock (= papillary stalk) that links the DP and the perifollicular CTS is thickened and now contains more than three cell layers (Fig 5F).

In pigmented HF, as in C57BL/6NCrBR mice, catagen II is very easy to distinguish as a dramatic reduction of the number of melanin granules is found in the precortical hair matrix (Fig 5D, E) (Slominski *et al.*, 1994). TUNEL staining, as an auxiliary method, reveals that up to four TUNEL⁺ keratinocytes are present in the hair bulb (Lindner *et al.*, 1997).

Catagen III HF (Fig 5G–I) are visibly shorter than catagen II HF, although the bulb and DP are still located close to the panniculus carnosus (Fig 5G). Although the follicle is shorter, the latter is due to the simultaneously declining overall skin thickness (Fig 1a, 2b). The key characteristic of this stage is the onion-like shape of the DP, a feature not found in any other stage (Fig 5G, I). At this time the bulb is smaller and encloses no more than half of the DP (Fig 5G). The characteristic onion-like shape of the DP and the open end of the bulb are particularly well visualized by the AP staining method (Fig 5G, I). In pigmented HF, no melanin granules are seen in the precortical hair matrix and the keratogenous region (Fig 5H, I). The presence of more than four TUNEL⁺ in the bulb is an auxiliary criterion for recognizing catagen III HF.

As for catagen III in catagen IV (Fig 5J–L) the shortening of the HF is compensated by the substantial, catagen-associated reduction in skin thickness so that the hair bulb and DP remain located close to the panniculus carnosus (Figs 4P and 5J). The key parameter for recognizing this stage is the compact, ball-like shape of the DP and the retracted bulb both clearly demarcated by the AP staining method (Fig 5K). In pigmented HF, only the distal part of the hair shaft is still pigmented (Fig 5J). Using NCAM immunoreactivity as an auxiliary marker (Müller-Röver *et al.*, 1998), only two to three trailing NCAM⁺ CTS cells are visible. Numerous TUNEL⁺ keratinocytes are now present in the bulb.

Recognition of catagen V–VIII HF Catagen V–VI HF (Fig 5M–R) display a characteristic “epithelial strand” between the secondary germ capsule and the DP. Whereas this epithelial strand represents only a small insertion between the DP and germ capsule in catagen V HF, catagen VI and VII HF display an epithelial strand that is substantially longer than the diameter of the DP (Fig 5P–U). Catagen VI–VIII HF can be differentiated by the length of the trailing CTS “tail” proximal to the DP: the trailing CTS tail of catagen VI HF is not longer than the DP diameter (Fig 5Q, R), whereas it is twice as long as the DP diameter during catagen VII (Fig 5S–U) and three times as long during catagen VIII (Fig 5V, X). Anti-NCAM labeling should be used as an auxiliary method in order to determine the precise length of the trailing NCAM⁺ CTS tail—one of the major parameters for distinguishing catagen VI, VII, and VIII, and telogen (Fig 5R, W) (Müller-Röver *et al.*, 1998).

During catagen V, the overall skin thickness as well as the HF length are substantially reduced. Yet, the compact, ball-shaped DP is still located in the subcutis (Fig 5M–O). The key characteristics of this stage are: (i) the development of the club hair, and (ii) the formation of the secondary germ capsule around the proximal end of the developing club hair (Fig 5M–O) leading to a constriction of the developing epithelial strand between the DP and germ capsule. The developing club hair base is characterized as a brush-like

structure at the most proximal depigmented end of the hair shaft, which is surrounded by about three cell layers of the developing secondary hair germ (ORS) (Fig 5M–O). Furthermore, the IRS begins to appear more opaque, and displays more sharply demarcated borders (Fig 5M). Using NCAM immunoreactivity as an auxiliary method reveals a trailing “tail” of NCAM⁺ CTS cells proximal to the DP, which may be as long as the diameter of the DP (Fig 5O).

Catagen VI HF (Fig 5P–R) are characterized by a clearly visible, brush-like hair club base that is entirely surrounded by the germ capsule (ORS) and has started to move distally (Fig 5Q). The opaque IRS is now very sharply demarcated from the ORS (Fig 5Q). The epithelial strand is longer than the diameter of the DP (Fig 5R). It is surrounded by a thick, glassy membrane, and contains multiple apoptotic (i.e., small pyknotic) keratinocytes that are frequently visible. The NCAM⁺ tail of trailing CTS fibroblasts is not longer than the diameter of the DP (Fig 5R).

The key parameter distinguishing catagen VII HF (Fig 5S–U) from catagen VI is the upward movement of the DP, leaving behind a long tail of CTS fibroblasts (Fig 5S–U). Stenn *et al.* (1998) proposed the concept of the “apoptotic force” to explain the upward movement of the DP (cf. Stenn and Paus, 2001): as catagen HF are characterized by an increased rate of apoptotic cell death in the bulb and epithelial strand (Lindner *et al.*, 1997) the volume of the proximal HF is increasingly reduced. Because the surviving keratinocytes are still linked by adhesion receptors such as E- and P-cadherin (Müller-Röver *et al.*, 1999), the progressively reduced size of the proximal HF moves the attached DP upwards within the CTS. The CTS tail is at least twice as long as the DP diameter (Fig 5S–U). Although the CTS tail is now clearly visible by light microscopy, it is particularly helpful to use anti-NCAM labeling to determine the precise length of the NCAM⁺ CTS tail, which is now more than twice as long as the DP diameter during catagen VII. The epithelial strand – still at least twice as long as the DP diameter (Fig 5S–U) – displays numerous apoptotic cells (Fig 5U) (Lindner *et al.*, 1997). AP labeling helps to locate the DP precisely, which has now moved upwards, resting in the middle of the subcutis (Fig 5T).

Catagen VIII HF (Fig 5V–X) are substantially shorter than catagen VII HF. Now the germ capsule as well as the club hair base reside in the dermis (Fig 5V, W). The DP is located at the border between the dermis and subcutis (Fig 5V, W). The trailing CTS tail, which consists of the folded glassy membrane as well as some fibroblasts and macrophages, still resides in the subcutis (Fig 5V, X). Remnants of the IRS are still present at the level of the SG (Fig 5V, X). In order to avoid the common error of mistaking the CTS tail or the secondary hair germ for the DP, it is again very helpful to use the AP staining technique.

Catagen VIII and telogen HF are distinguished by the length of the trailing CTS using NCAM immunoreactivity: the trailing NCAM⁺ CTS is more than three times longer than the DP diameter in catagen VIII HF (Fig 5W), whereas it is absent in telogen HF.

ANALYSIS OF HAIR CYCLE ALTERATIONS IN MUTANT OR PHARMACOLOGICALLY TREATED MICE

Using the qualitative criteria characterized in this guide, it is possible to provide fully quantitative comparisons of abnormalities in HF cycling in mutant (transgenic, knockout, spontaneous mutants) or pharmacologically treated mice compared with age-matched wild-type or vehicle controls. For quantitative experiments, a sufficiently large number of age- and sex-matched test and control mice should be compared by quantitative histomorphometry, i.e., the morphologic criteria described above should be used to determine the stages of a defined number of longitudinally cut HF per mouse (e.g., 20 HF per mouse, studying three to five mice per time point and group). Quantitative histomorphometry for the study of abnormalities in murine HF cycling, using the classification criteria suggested here, has been used by us in previous

publications (Paus *et al*, 1994b; Maurer *et al*, 1997b, 1998; Schilli *et al*, 1997; Botchkarev *et al*, 1998, 1999; Müller-Röver *et al*, 2000a, b).

For the qualitative and quantitative evaluation of HF it is of pivotal importance to study standardized longitudinal sections of HF, prepared by using a special harvesting and embedding technique (Paus *et al*, 1999). A sufficient number of well-cut skin sections has to be prepared for each mouse in order to assign the HF to defined hair cycle stages. The total number of HF per hair cycle stage (anagen I–VI, catagen I–VIII, telogen) then can be compared quantitatively and statistically between test and control mice.

This quantitative approach can be developed further by calculating a "hair cycle score" as described previously (Maurer *et al*, 1997a, b). This allows one to compare either anagen or catagen development between large cohorts of HF from test and control groups of mice. To this end every stage of anagen or catagen is assigned a factor in ascending numerical order (e.g. for anagen: anagen I = factor 1, anagen II = factor 2, anagen III = factor 3, etc.; for catagen: catagen I = factor 1, catagen II = factor 2, catagen III = factor 3, etc.). The number of HF in each specific stage is multiplied by the corresponding factor. The results of each sum are totalled and divided by the total number of HF counted. This gives a final value between 1 and 6 for anagen HF and between 1 and 8 for catagen HF, thus defining the average stage of all HF within the entire group. This will allow one to identify even subtle abnormalities in the dynamics of HF cycling between test and control mice that might otherwise have escaped notice. In addition, we have to highlight that HF located close to each other very rarely display vastly different stages; therefore, "outlying" HF stages are also very rarely seen. In these rare cases, however, the statistical results have to be interpreted carefully.

More severe alterations of the skin phenotype such as might be caused by a spontaneous or experimentally induced mutation, can result in a large number of alterations to the HF phenotype (Sundberg, 1994), including the induction of HF dystrophy, i.e., morphologic signs of HF damage such as abnormally distended hair canals, massive apoptosis, and ectopic melanin granules (cf. Paus *et al*, 1994c, 1996; Maurer *et al*, 1997b; Schilli *et al*, 1998). A drug effect on hair cycling or the pathogenesis of a mutation can only be understood fully if the exact stage is known when the aberrations are seen for the first time (for a good example see Panteleyev *et al*, 1998). In any case, critical analysis of alterations in the speed and synchronization of HF cycling, and in the morphology of defined HF compartments requires implementation and quantification of such assessment criteria as suggested in this guide.

In summary, this review offers a comprehensive, pragmatic, and simple guide to recognizing the morphologic features of murine HF cycling, and serves as a useful tool for rapid and instructive analyses of mouse hair phenotypes after genetic or pharmacologic manipulations. The schematic drawings of individual hair cycle stages presented here may be employed by other investigators for reporting the follicular expression patterns of their genes and proteins of interest so that the documentation of such expression patterns will provide a framework for a standardized and highly reproducible comparison.

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